SEQUENCE LISTING

10

In the following SEQ ID Nos. 1, 3, 5 the 5', coding sequence and 3' sequence of the relevant α -amylase genes are illustrated. The 5' sequence is the first separate part of the sequence written with lower case letters, the coding sequence is the intermediate part of the sequence, where the signal sequence is written with lower case letters and the sequence encoding the mature α -amylase is written with upper case letters, and the 3' sequence is the third separate part of the sequence written with lower case letters.

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<211> 1814

<212> DNA

<213> Bacillus stearothermohilus

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                                                                       120
                                                                       180
tcgaatgtaa catttgatta agggggaagg gcattgtgct aacgtttcac cgcatcattc
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                                                                       300
gacagecege caaggetgee geacegttta aeggeaceat gatgeagtat tttgaatggt
                                                                       360
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                                                                       420
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tagggtacgg agtatacgac ttgtatgacc tcggcgaatt caatcaaaaa gggaccgtcc
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                                                                       540
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gaatgcaagt gtacgccgat gtcgtgttcg accataaagg cggcgctgac ggcacggaat
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                                                                       660
                                                                       720
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actatgacta cttaatgtat geegacettg atatggatea teeegaagte gtgacegage
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                                                                       960
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ctggcaagcc gctatttacc gtcggggaat attggagcta tgacatcaac aagttgcaca
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                                                                      1260
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ataacattcc ttcgctgaaa agcaaaatcg atccgctcct catcgcgcgc agggattatg
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                                                                      1680
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gttcggtttc ggtttgggtt cctagaaaaa cgaccgtttc taccatcgct cggccgatca
                                                                      1740
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<210> 6 <211> 514 <212> PRT

<213> Bacillus stearothermophilus

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			180					185					190		
Asn	Gly	Asn 195	Tyr	Asp	Tyr	Leu	Met 200	Tyr	Ala	Asp	Leu	Asp 205	Met	Asp	His
Pro	Glu 210	Val	Val	Thr	Glu	Leu 215	Lys	Asn	Trp	Gly	Lys 220	Trp	Tyr	Val	Asn
225					230					235			His		240
Phe	Ser	Phe	Phe	Pro 245	Asp	Trp	Leu	Ser	Tyr 250	Val	Arg	Ser	Gln	Thr 255	Gly
Lys	Pro	Leu	Phe 260	Thr	Val	Gly	Glu	Tyr 265	Trp	Ser	Tyr	Asp	Ile 270	Asn	Lys
Leu	His	Asn 275	Tyr	Ile	Thr	Lys	Thr 280	Asp	Gly	Thr	Met	Ser 285	Leu	Phe	Asp
Ala	Pro	Leu	His	Asn	Lys	Phe 295	Tyr	Thr	Ala	Ser	Lys 300	Ser	Gly	Gly	Ala
Phe	Asp	Met	Arg	Thr	Leu 310	Met	Thr	Asn	Thr	Leu 315	Met	Lys	Asp	Gln	Pro 320
	Leu	Ala	Val	Thr	Phe	Val	Asp	Asn	His	Asp	Thr	Glu	Pro	Gly 335	Gln

Ala Leu Gln Ser Trp Val Asp Pro Trp Phe Lys Pro Leu Ala Tyr Ala Phe Ile Leu Thr Arg Gln Glu Gly Tyr Pro Cys Val Phe Tyr Gly Asp Tyr Tyr Gly Ile Pro Gln Tyr Asn Ile Pro Ser Leu Lys Ser Lys Ile Asp Pro Leu Leu Ile Ala Arg Arg Asp Tyr Ala Tyr Gly Thr Gln His Asp Tyr Leu Asp His Ser Asp Ile Ile Gly Trp Thr Arg Glu Gly Gly Thr Glu Lys Pro Gly Ser Gly Leu Ala Ala Leu Ile Thr Asp Gly Pro Gly Gly Ser Lys Trp Met Tyr Val Gly Lys Gln His Ala Gly Lys Val Phe Tyr Asp Leu Thr Gly Asn Arg Ser Asp Thr Val Thr Ile Asn Ser Asp Gly Trp Gly Glu Phe Lys Val Asn Gly Gly Ser Val Ser Val Trp Val Pro Arg Lys Thr Thr Val Ser Thr Ile Ala Arg Pro Ile Thr Thr Arg Pro Trp Thr Gly Glu Phe Val Arg Trp Thr Glu Pro Arg Leu Val Ala Trp

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<213> Bacillus licheniformis

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Pro Leu Leu Asn Ala Phe Lys Ser Thr Ser Gly Ser Met Asp Asp Leu
                               265
Tyr Asn Met Ile Asn Thr Val Lys Ser Asp Cys Pro Asp Ser Thr Leu
                            280
                                                285
        275
Leu Gly Thr Phe Val Glu Asn His Asp Asn Pro Arg Phe Ala Ser Tyr
    290
                        295
                                            300
Thr Asn Asp Ile Ala Leu Ala Lys Asn Val Ala Ala Phe Ile Ile Leu
                    310
                                        315
Asn Asp Gly Ile Pro Ile Ile Tyr Ala Gly Gln Glu Gln His Tyr Ala
                325
                                    330
                                                        335
Gly Gly Asn Asp Pro Ala Asn Arg Glu Ala Thr Trp Leu Ser Gly Tyr
                                345
Pro Thr Asp Ser Glu Leu Tyr Lys Leu Ile Ala Ser Ala Asn Ala Ile
                          360
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Arg Asn Tyr Ala Ile Ser Lys Asp Thr Gly Phe Val Thr Tyr Lys Asn
                       375
                                           380
Trp Pro Ile Tyr Lys Asp Asp Ile Thr Ile Ala Met Arg Lys Gly Thr
                    390
                                        395
Asp Gly Ser Gln Ile Val Thr Ile Leu Ser Asn Lys Gly Ala Ser Gly
                405
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Asp Ser Tyr Thr Leu Ser Leu Ser Gly Ala Gly Tyr Thr Ala Gly Gln
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Gln Leu Thr Glu Val Ile Gly Cys Thr Thr Val Thr Val Gly Ser Asp
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Gly Asn Val Pro Val Pro Met Ala Gly Gly Leu Pro Arg Val Leu Tyr
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Pro Thr Glu Lys Leu Ala Gly Ser Lys Ile Cys Ser Ser Ser
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      <210> 11
      <211> 36
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	l2> DNA l3> Bacillus licheniformis	
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	l0> 17 l1> 30	
	l2> DNA l3> Bacillus licheniformis	
	00> 17 cc gaattcattg gaaactteec	30
	lO> 18 l1> 34	
	12> DNA 13> Bacillus licheniformis	
-	00> 18 cc gaattcaggg gaaacttccc aatc	34
	lO> 19 l1> 41	
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      <400> 29
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                                                                         75
qacqagtccc gaaag
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      <211> 84
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                                                                         60
                                                                         84
rskaactatg attatttgat gtat
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tggtatgcca at
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      <223> N at 31 is 93% C, 2% T, 2% A, 3% G
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gttaatcatg tcagggaa
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      <211> 63
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                                                                         63
ttt
      <210> 36
      <211> 78
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                                                                         78
tttcaaggaa aggcttgg
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      <211> 93
      <212> DNA
      <213 > Bacillus licheniformis
      <220>
      <221> misc_feature
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      <400> 37
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                                                                         60
                                                                         93
avcargacar attttaatca ttcagtgttt gac
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